

34 to the number of amino acids (AA) counted after the AA at the start of the N terminus in VP-1 of AAV2). In the subsequent PCR there is use of in each case 2 mutation-specific primers and, as template, a plasmid, pCap, which contains only the cap gene and is formed by cutting out the 2.2 kb

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In the Claims:

Amend claim 14 as follows.

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35 14. (Twice Amended) Structural protein according to Claim 8, characterized in that one or more insertions in VP3 is/are located before and/or after at least one amino acid in the sequence selected from YKQIS SQSGA (SEQ ID NO: 2), YLTLN NGSQA (SEQ ID NO: 3), YYLSR TNTPS (SEQ ID NO: 4), EEKFF PQSGV (SEQ ID NO: 5), NPVAT, EQYGS (SEQ ID NO: 6), LQRGN RQAAT (SEQ ID NO: 7), and NVDFTVDTNG (SEQ ID NO: 8).

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